SEQUENCE LISTING

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Ramsdell, Fred

<120> IDENTIFICATION OF THE GENE CAUSING THE
MOUSE SCURFY PHENOTYPE AND ITS HUMAN ORTHOLOG

<130> 240083.501D6

<140> US 09/696,867
<141> 2000-10-25

<160> 14

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<210> 1
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<213> Mus musculus

<400> 1

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        35                  40                  45
Asp Leu Arg Ser Gly Ala His Thr Ser Ser Ser Leu Asn Pro Leu Pro
    50                  55                  60
Pro Ser Gln Leu Gln Leu Pro Thr Val Pro Leu Val Met Val Ala Pro
65                  70                  75                  80
Ser Gly Ala Arg Leu Gly Pro Ser Pro His Leu Gln Ala Leu Leu Gln
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Asp Arg Pro His Phe Met His Gln Leu Ser Thr Val Asp Ala His Ala
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Gln Thr Pro Val Leu Gln Val Arg Pro Leu Asp Asn Pro Ala Met Ile
        115                 120                 125
Ser Leu Pro Pro Pro Ser Ala Ala Thr Gly Val Phe Ser Leu Lys Ala
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Arg Pro Gly Leu Pro Pro Gly Ile Asn Val Ala Ser Leu Glu Trp Val
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Ser Arg Glu Pro Ala Leu Leu Cys Thr Phe Pro Arg Ser Gly Thr Pro
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Arg Lys Asp Ser Asn Leu Leu Ala Ala Pro Gln Gly Ser Tyr Pro Leu
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Leu Ala Asn Gly Val Cys Lys Trp Pro Gly Cys Glu Lys Val Phe Glu
        195                 200                 205
Glu Pro Glu Glu Phe Leu Lys His Cys Gln Ala Asp His Leu Leu Asp
    210                 215                 220
Glu Lys Gly Lys Ala Gln Cys Leu Leu Gln Arg Glu Val Val Gln Ser
225                 230                 235                 240
Leu Glu Gln Gln Leu Glu Leu Glu Lys Glu Lys Leu Gly Ala Met Gln
                245                 250                 255
Ala His Leu Ala Gly Lys Met Ala Leu Ala Lys Ala Pro Ser Val Ala
            260                 265                 270
Ser Met Asp Lys Ser Ser Cys Cys Ile Val Ala Thr Ser Thr Gln Gly
        275                 280                 285
Ser Val Leu Pro Ala Trp Ser Ala Pro Arg Glu Ala Pro Asp Gly Gly
    290                 295                 300
Leu Phe Ala Val Arg Arg His Leu Trp Gly Ser His Gly Asn Ser Ser
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Phe Pro Glu Phe Phe His Asn Met Asp Tyr Phe Lys Tyr His Asn Met
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Ala Pro Glu Arg Gln Arg Thr Leu Asn Glu Ile Tyr His Trp Phe Thr
            355                   360                   365
Arg Met Phe Ala Tyr Phe Arg Asn His Pro Ala Thr Trp Lys Asn Ala
        370                   375                   380
Ile Arg His Asn Leu Ser Leu His Lys Cys Phe Val Arg Val Glu Ser
385                   390                   395                   400
Glu Lys Gly Ala Val Trp Thr Val Asp Glu Phe Glu Phe Arg Lys Lys
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Arg Ser Gln Arg Pro Asn Lys Cys Ser Asn Pro Cys Pro
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<211> 1869
<212> DNA
<213> Homo sapien

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tcccacaagc caggctgatc cttttctgtc agtccacttc accaagcctg cccttggaca    180
aggacccgat gcccaacccc aggcctggca agccctcggc cccttccttg gcccttggcc    240
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cccggggccc aggggggaacc ttccagggcc gagatcttcg aggcggggcc catgcctcct    360
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tgcaggtgca ccccctggag agcccagcca tgatcagcct cacaccaccc accaccgcca    600
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<212> PRT
<213> Homo sapien
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<400> 4
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            20              25              30
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        35              40              45
Asp Leu Arg Gly Gly Ala His Ala Ser Ser Ser Ser Leu Asn Pro Met
    50              55              60
Pro Pro Ser Gln Leu Gln Leu Pro Thr Leu Pro Leu Val Met Val Ala
65              70              75              80
Pro Ser Gly Ala Arg Leu Gly Pro Leu Pro His Leu Gln Ala Leu Leu
            85              90              95
Gln Asp Arg Pro His Phe Met His Gln Leu Ser Thr Val Asp Ala His
            100             105             110
Ala Arg Thr Pro Val Leu Gln Val His Pro Leu Glu Ser Pro Ala Met
            115             120             125
Ile Ser Leu Thr Pro Pro Thr Thr Ala Thr Gly Val Phe Ser Leu Lys
    130             135             140
Ala Arg Pro Gly Leu Pro Pro Gly Ile Asn Val Ala Ser Leu Glu Trp
145             150             155             160
Val Ser Arg Glu Pro Ala Leu Leu Cys Thr Phe Pro Asn Pro Ser Ala
            165             170             175
Pro Arg Lys Asp Ser Thr Leu Ser Ala Val Pro Gln Ser Ser Tyr Pro
            180             185             190
Leu Leu Ala Asn Gly Val Cys Lys Trp Pro Gly Cys Glu Lys Val Phe
            195             200             205
Glu Glu Pro Glu Asp Phe Leu Lys His Cys Gln Ala Asp His Leu Leu
    210             215             220
Asp Glu Lys Gly Arg Ala Gln Cys Leu Leu Gln Arg Glu Met Val Gln
225             230             235             240
Ser Leu Glu Gln Gln Leu Val Leu Glu Lys Glu Lys Leu Ser Ala Met
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Gln Ala His Leu Ala Gly Lys Met Ala Leu Thr Lys Ala Ser Ser Val
            260             265             270
Ala Ser Ser Asp Lys Gly Ser Cys Cys Ile Val Ala Ala Gly Ser Gln
    275             280             285
Gly Pro Val Val Pro Ala Trp Ser Gly Pro Arg Glu Ala Pro Asp Ser
    290             295             300
Leu Phe Ala Val Arg Arg His Leu Trp Gly Ser His Gly Asn Ser Thr
305             310             315             320
Phe Pro Glu Phe Leu His Asn Met Asp Tyr Phe Lys Phe His Asn Met
            325             330             335
Arg Pro Pro Phe Thr Tyr Ala Thr Leu Ile Arg Trp Ala Ile Leu Glu
            340             345             350
Ala Pro Glu Lys Gln Arg Thr Leu Asn Glu Ile Tyr His Trp Phe Thr
            355             360             365
Arg Met Phe Ala Phe Phe Arg Asn His Pro Ala Thr Trp Lys Asn Ala
    370             375             380
Ile Arg His Asn Leu Ser Leu His Lys Cys Phe Val Arg Val Glu Ser
385             390             395             400
Glu Lys Gly Ala Val Trp Thr Val Asp Glu Leu Glu Phe Arg Lys Lys
            405             410             415
Arg Ser Gln Arg Pro Ser Arg Cys Ser Asn Pro Thr Pro Gly Pro
            420             425             430

<210> 5
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<211> 23
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<213> Artificial Sequence

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<223> Primer for generation of mouse Fkh cDNA

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<400> 7
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<210> 10
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<213> Artificial Sequence

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<223> Primer for PCR amplification of mouse Fkh cDNA

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<210> 11
<211> 28
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<223> Oligonucleotide for hybridization reaction

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<210> 12
<211> 28
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<220>
<223> Oligonucleotide for hybridization reaction

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<223> n = C or T

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<223> n = G or A

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<210> 14
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<223> n = C or T

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<223> n = G or A

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